

# Class17

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```
vax <- read.csv("covid19vaccinesbyzipcode_test.csv")
head(vax)
```

```
##   as_of_date zip_code_tabulation_area local_health_jurisdiction      county
## 1 2021-01-05           92395           San Bernardino San Bernardino
## 2 2021-01-05           93206                Kern           Kern
## 3 2021-01-05           91006           Los Angeles Los Angeles
## 4 2021-01-05           91901           San Diego San Diego
## 5 2021-01-05           92230           Riverside Riverside
## 6 2021-01-05           92662            Orange      Orange
##   vaccine_equity_metric_quartile      vem_source
## 1                             1 Healthy Places Index Score
## 2                             1 Healthy Places Index Score
## 3                             3 Healthy Places Index Score
## 4                             3 Healthy Places Index Score
## 5                             1 Healthy Places Index Score
## 6                             4 Healthy Places Index Score
##   age12_plus_population age5_plus_population persons_fully_vaccinated
## 1                35915.3                40888                NA
## 2                 1237.5                 1521                NA
## 3                28742.7                31347                19
## 4                15549.8                16905                12
## 5                 2320.2                 2526                NA
## 6                 2349.5                 2397                NA
##   persons_partially_vaccinated percent_of_population_fully_vaccinated
## 1                        NA                        NA
## 2                        NA                        NA
## 3                        873                        0.000606
## 4                        271                        0.000710
## 5                        NA                        NA
## 6                        NA                        NA
##   percent_of_population_partially_vaccinated
## 1                        NA
## 2                        NA
## 3                        0.027850
## 4                        0.016031
## 5                        NA
## 6                        NA
##   percent_of_population_with_1_plus_dose
## 1                        NA
## 2                        NA
```

```
## 3 0.028456
## 4 0.016741
## 5 NA
## 6 NA
##
## 1 Information redacted in accordance with CA state privacy requirements
## 2 Information redacted in accordance with CA state privacy requirements
## 3 No
## 4 No
## 5 Information redacted in accordance with CA state privacy requirements
## 6 Information redacted in accordance with CA state privacy requirements
```

```
nrow(vax)
```

```
## [1] 82908
```

```
skimr::skim(vax)
```

Table 1: Data summary

Name	vax
Number of rows	82908
Number of columns	14
Column type frequency:	
character	5
numeric	9
Group variables	None

### Variable type: character

skim_variable	n_missing	complete_rate	min	max	empty	n_unique	whitespace
as_of_date	0	1	10	10	0	47	0
local_health_jurisdiction	0	1	0	15	235	62	0
county	0	1	0	15	235	59	0
vem_source	0	1	15	26	0	3	0
redacted	0	1	2	69	0	2	0

### Variable type: numeric

skim_variable	n_missing	complete_rate	mean	sd	p0	p25	p50	p75	p100	hist
zip_code_tabulation_area	0	1.00	93665.18	17.39	00001	192257.75	3658.55	380.57	635.0	
vaccine_equity_metric_4qartile	0	0.95	2.44	1.11	1	1.00	2.00	3.00	4.0	
age12_plus_population	0	1.00	18895.04	993.94	0	1346.95	3685.11	756.18	556.7	
age5_plus_population	0	1.00	20875.21	1106.04	0	1460.50	5364.01	877.00	1902.0	
persons_fully_vaccinated	8355	0.90	9585.35	1609.12	1	516.00	4210.00	6095.00	1219.0	
persons_partially_vaccinated	8855	0.90	1894.87	2105.55	11	198.00	1269.00	2880.00	20159.0	
percent_of_population_fully_vaccinated	8355	0.90	0.43	0.27	0	0.20	0.44	0.63	1.0	

skim_variable	n_missing	complete	mean	sd	p0	p25	p50	p75	p100	hist
percent_of_population_855	0	100	0.99	0.10	0	0.06	0.07	0.11	1.0	
percent_of_population_855h_1_000s_0051	0	100	0.51	0.26	0	0.31	0.53	0.71	1.0	

```
library(lubridate)
```

```
## Warning: package 'lubridate' was built under R version 4.1.2
```

```
##
```

```
## Attaching package: 'lubridate'
```

```
## The following objects are masked from 'package:base':
```

```
##
```

```
## date, intersect, setdiff, union
```

```
today()
```

```
## [1] "2021-11-24"
```

```
vax$as_of_date[1]
```

```
## [1] "2021-01-05"
```

```
vax$as_of_date <- ymd(vax$as_of_date)
```

```
today() - vax$as_of_date[1]
```

```
## Time difference of 323 days
```

```
today() - vax$as_of_date[nrow(vax)]
```

```
## Time difference of 1 days
```

```
vax$as_of_date[nrow(vax)] - vax$as_of_date[1]
```

```
## Time difference of 322 days
```

```
length(unique((vax$zip_code_tabulation_area)))
```

```
## [1] 1764
```

```
library(zipcodeR)
```

```
## Warning: package 'zipcodeR' was built under R version 4.1.2
```

```
geocode_zip('92037')
```

```
## # A tibble: 1 x 3
##   zipcode lat lng
##   <chr>   <dbl> <dbl>
## 1 92037   32.8 -117.
```

```
zip_distance('92037','92109')
```

```
##   zipcode_a zipcode_b distance
## 1      92037      92109     2.33
```

```
reverse_zipcode(c('92037', "92109") )
```

```
## # A tibble: 2 x 24
##   zipcode zipcode_type major_city post_office_city common_city_list county state
##   <chr>   <chr>         <chr>         <chr>         <blob> <chr> <chr>
## 1 92037   Standard      La Jolla     La Jolla, CA      <raw 20 B> San D~ CA
## 2 92109   Standard      San Diego    San Diego, CA      <raw 21 B> San D~ CA
## # ... with 17 more variables: lat <dbl>, lng <dbl>, timezone <chr>,
## #   radius_in_miles <dbl>, area_code_list <blob>, population <int>,
## #   population_density <dbl>, land_area_in_sqmi <dbl>,
## #   water_area_in_sqmi <dbl>, housing_units <int>,
## #   occupied_housing_units <int>, median_home_value <int>,
## #   median_household_income <int>, bounds_west <dbl>, bounds_east <dbl>,
## #   bounds_north <dbl>, bounds_south <dbl>
```

```
library(dplyr)
```

```
##
## Attaching package: 'dplyr'

## The following objects are masked from 'package:stats':
##
##   filter, lag

## The following objects are masked from 'package:base':
##
##   intersect, setdiff, setequal, union
```

```
inds <- vax$county == "San Diego"
nrow(vax[inds,])
```

```
## [1] 5029
```

```
sd <- filter(vax, county == "San Diego")
nrow(sd)
```

```
## [1] 5029
```

```
sd.20 <- filter(vax, county == "San Diego" &
  age5_plus_population > 20000)
nrow(sd.20)
```

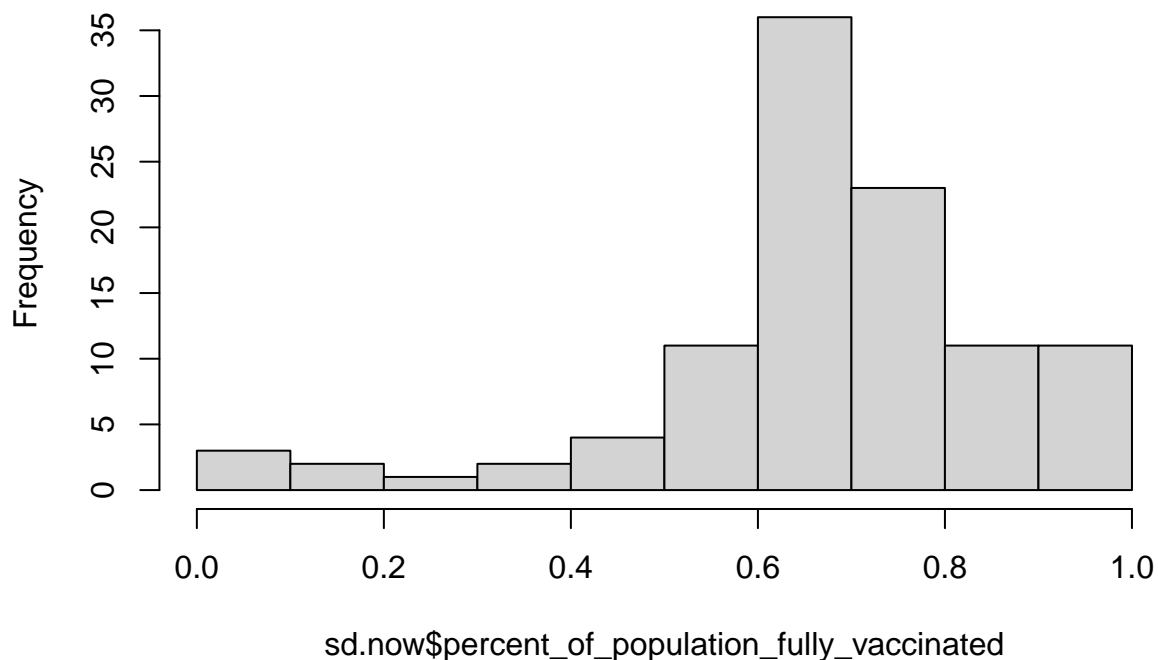
```
## [1] 3055
```

```
sd.now <- filter(sd, as_of_date == "2021-11-23")
summary(sd.now$percent_of_population_fully_vaccinated)
```

```
##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.     NA's
## 0.01017 0.61301 0.67965 0.67400 0.76932 1.00000         3
```

```
hist(sd.now$percent_of_population_fully_vaccinated)
```

## Histogram of sd.now\$percent\_of\_population\_fully\_vaccinated



```
head(sd.now)
```

```
##   as_of_date zip_code_tabulation_area local_health_jurisdiction  county
## 1 2021-11-23           92120                San Diego San Diego
## 2 2021-11-23           91962                San Diego San Diego
## 3 2021-11-23           92155                San Diego San Diego
## 4 2021-11-23           92147                San Diego San Diego
## 5 2021-11-23           91913                San Diego San Diego
## 6 2021-11-23           92114                San Diego San Diego
```

```
## vaccine_equity_metric_quartile vem_source
## 1 4 Healthy Places Index Score
## 2 3 Healthy Places Index Score
## 3 NA No VEM Assigned
## 4 NA No VEM Assigned
## 5 3 Healthy Places Index Score
## 6 2 Healthy Places Index Score
## age12_plus_population age5_plus_population persons_fully_vaccinated
## 1 26372.9 28414 21234
## 2 1758.7 2020 948
## 3 456.0 456 70
## 4 518.0 518 NA
## 5 43514.7 50461 37974
## 6 59050.7 64945 43708
## persons_partially_vaccinated percent_of_population_fully_vaccinated
## 1 3198 0.747308
## 2 126 0.469307
## 3 20 0.153509
## 4 NA NA
## 5 6690 0.752542
## 6 6261 0.673000
## percent_of_population_partially_vaccinated
## 1 0.112550
## 2 0.062376
## 3 0.043860
## 4 NA
## 5 0.132578
## 6 0.096405
## percent_of_population_with_1_plus_dose
## 1 0.859858
## 2 0.531683
## 3 0.197369
## 4 NA
## 5 0.885120
## 6 0.769405
## redacted
## 1 No
## 2 No
## 3 No
## 4 Information redacted in accordance with CA state privacy requirements
## 5 No
## 6 No
```

```
lj <- filter(sd.now, zip_code_tabulation_area == "92037")
```

```
lj$age5_plus_population
```

```
## [1] 36144
```

```
lj$percent_of_population_fully_vaccinated
```

```
## [1] 0.916196
```

```
lj2 <- filter(sd.now, zip_code_tabulation_area == "92122")
lj2$age5_plus_population
```

```
## [1] 45951
```

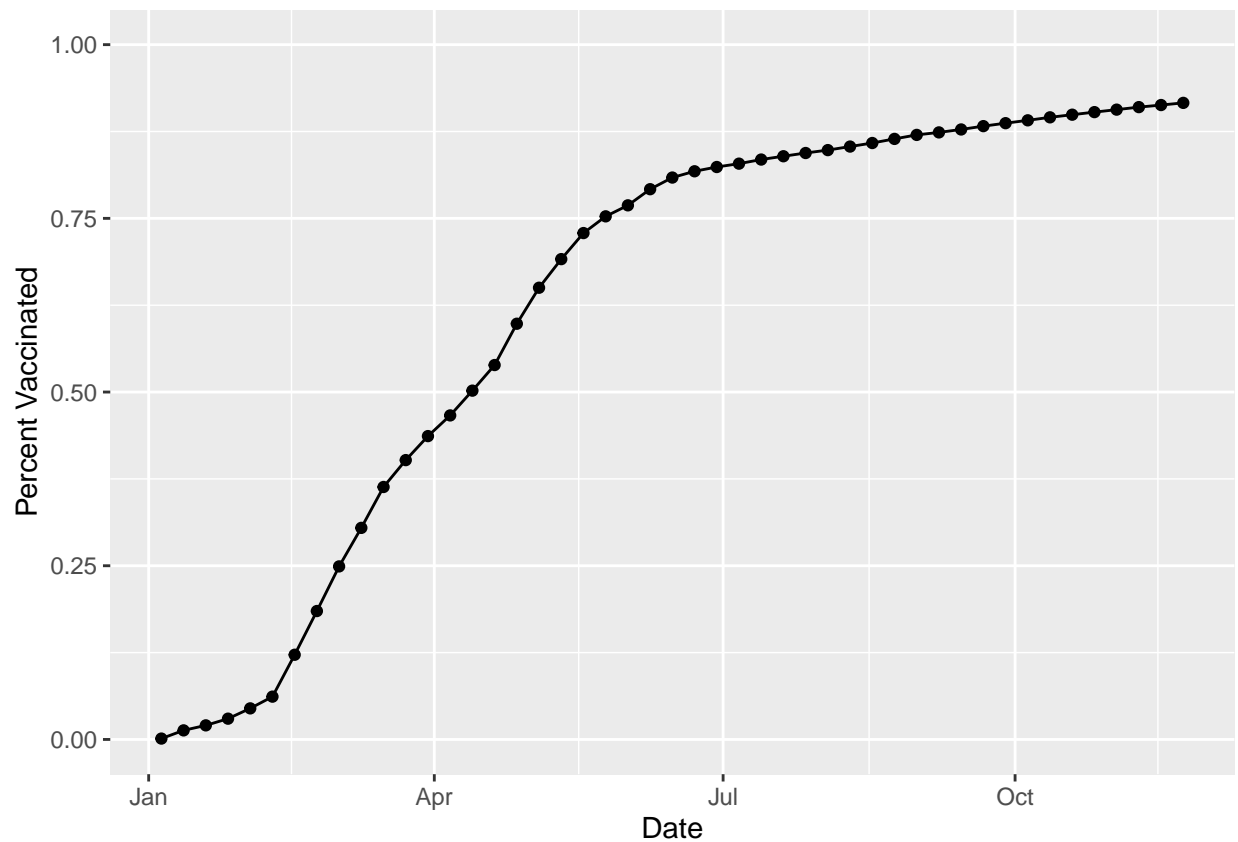
```
lj2$percent_of_population_fully_vaccinated
```

```
## [1] 0.771474
```

```
library(ggplot2)
```

```
lj3 <- filter(vax, zip_code_tabulation_area == "92037")
```

```
ggplot(lj3) +
  aes(x=as_of_date,
      y=percent_of_population_fully_vaccinated) +
  geom_point() +
  geom_line(group=1) + ylim(c(0,1)) +
  labs(x= "Date", y="Percent Vaccinated")
```



```
sd.36 <- filter(vax, county== "San Diego", age5_plus_population>36144 )
head(sd.36)
```

```

##   as_of_date zip_code_tabulation_area local_health_jurisdiction   county
## 1 2021-01-05                92058             San Diego San Diego
## 2 2021-01-05                92078             San Diego San Diego
## 3 2021-01-05                92019             San Diego San Diego
## 4 2021-01-05                92117             San Diego San Diego
## 5 2021-01-05                92057             San Diego San Diego
## 6 2021-01-05                91913             San Diego San Diego
##   vaccine_equity_metric_quartile          vem_source
## 1                             1 Healthy Places Index Score
## 2                             3 Healthy Places Index Score
## 3                             3 Healthy Places Index Score
## 4                             3 Healthy Places Index Score
## 5                             2 Healthy Places Index Score
## 6                             3 Healthy Places Index Score
##   age12_plus_population age5_plus_population persons_fully_vaccinated
## 1                34956.0                39695                NA
## 2                41789.5                47476                37
## 3                37439.4                40464                25
## 4                50041.6                53839                42
## 5                51927.0                56906                22
## 6                43514.7                50461                37
##   persons_partially_vaccinated percent_of_population_fully_vaccinated
## 1                        NA                        NA
## 2                        688                        0.000779
## 3                        610                        0.000618
## 4                       1143                        0.000780
## 5                        691                        0.000387
## 6                       1993                        0.000733
##   percent_of_population_partially_vaccinated
## 1                        NA
## 2                        0.014492
## 3                        0.015075
## 4                        0.021230
## 5                        0.012143
## 6                        0.039496
##   percent_of_population_with_1_plus_dose
## 1                        NA
## 2                        0.015271
## 3                        0.015693
## 4                        0.022010
## 5                        0.012530
## 6                        0.040229
##                                     redacted
## 1 Information redacted in accordance with CA state privacy requirements
## 2                                     No
## 3                                     No
## 4                                     No
## 5                                     No
## 6                                     No

```

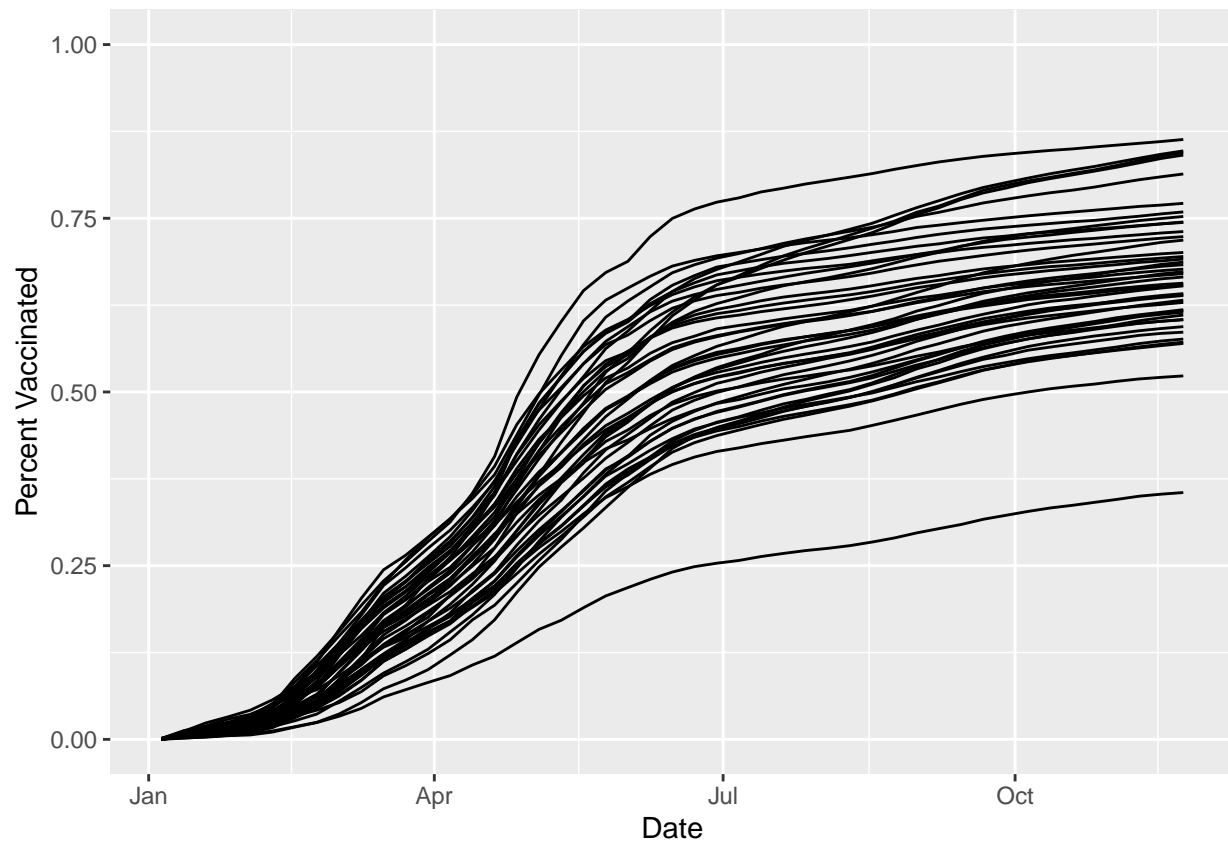
```
length(unique(sd.36$zip_code_tabulation_area))
```

```
## [1] 43
```



```
ggplot(sd.36) +
  aes(x=as_of_date,
      y=percent_of_population_fully_vaccinated, group=zip_code_tabulation_area) +
  geom_line() + ylim(c(0,1)) +
  labs(x= "Date", y="Percent Vaccinated")
```

```
## Warning: Removed 1 row(s) containing missing values (geom_path).
```

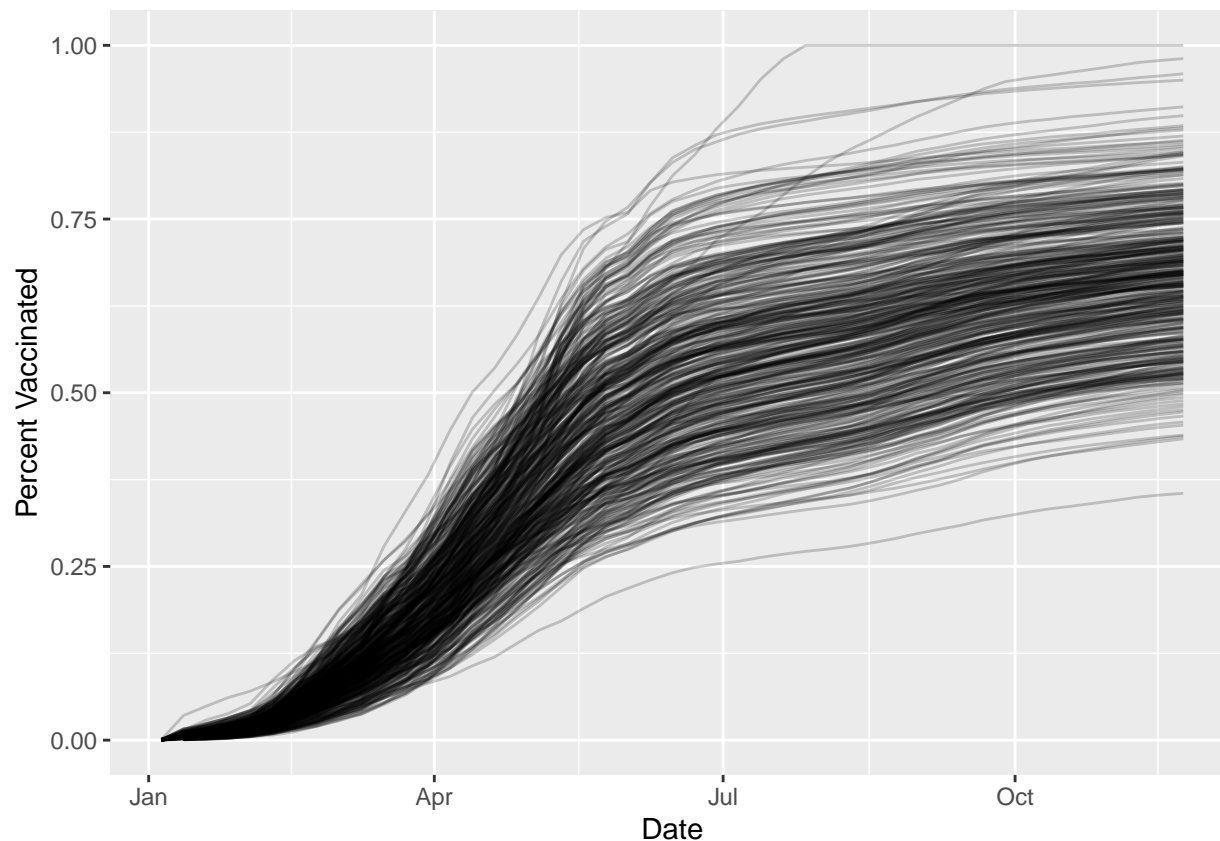


```
ca<- filter(vax, age5_plus_population>36144 )
length(unique(ca$zip_code_tabulation_area))
```

```
## [1] 411
```

```
ggplot(ca) +
  aes(x=as_of_date,
      y=percent_of_population_fully_vaccinated, group=zip_code_tabulation_area) +
  geom_line(alpha=0.2) + ylim(c(0,1)) +
  labs(x= "Date", y="Percent Vaccinated")
```

```
## Warning: Removed 176 row(s) containing missing values (geom_path).
```



```
ca.now <- filter(ca, as_of_date== "2021-11-23")
summary(ca.now$percent_of_population_fully_vaccinated)
```

```
##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
## 0.3552  0.5939  0.6696  0.6672  0.7338  1.0000
```

```
ggplot(ca) +
  aes(x=as_of_date,
      y=percent_of_population_fully_vaccinated, group=zip_code_tabulation_area) +
  geom_line(alpha=0.2) + ylim(c(0,1)) +
  labs(x= "Date", y="Percent Vaccinated") +
  geom_hline(yintercept = mean(ca.now$percent_of_population_fully_vaccinated), linetype= 2, color="red")
```

```
## Warning: Removed 176 row(s) containing missing values (geom_path).
```

